NE	B 1233							
	-1391	-1381	-1371	-1361 *	-1351 *	-1341	-1331 *	-1321
	AGCOGATAAC	AATTICACAC	<u>AGGA</u> AACAGC		ATTACGCCAA	GCTTGGTACC	GAGCTOOGAT -Bamil	
		To	c promoter_					
			c promocer_					
	-1311	-1301	-1291	-1281	-1271	-1261	-1251	-1241
	*	*	1.*			*	*	*
	COCCCCCAC	A CONTROLLAR			COSTGGTOGA	cccccccccc	TOOTAACTIT	AAGAGAAATT
		pCR2.1			0001001001	-SmaI-		
		pure.I		Ger	omeWalker A	Adaptor	>	
			_		20110110121102			mACO1>
	-1231	-1221	-1211	_1201	-1191	-1181	-1171	-1161
	-1231	-1221	-1211	-1201		*	*	*
	COURS & STREET	CONCRETE ON TO	מ מיניים מיניים מ	татассасаа	יים ביייייים בייי	TCTAATGTTG	ייידיייביינייניביי	CGATAAAGTT
	GGIAMMIIC					ichimorio		
				direct Ge	AMILE IAM			
	-1151	-1141	-1131	-1121	1111	-1101	-1091	-1081
	-1121	-1141	-1131	-1757	-1111	-1101	-1031	*
		macent or our	COMPONED CO	ma amercamini	CONTRACTOR	AATTTGTATC	TO ACIDITETE A	CHICADATIC
	AAATAAAGIG	TOGTAGACGA				AMITIGIAIC		GIICAMAIIC
				CITALOI GE	IOUTE DIVA			
		40.53	4054	1041	1001	-1021	-1011	-1001
	-1071	-1061	-1021	-1041		-1021	-1011	-1001
	*	*					***************************************	* * CUIDOTTA *C
	ACACTAAAAC					GATGTATICC		
				cmacor Ge	nomic Liva			
			054	-961	-951	-941	-931	-921
	-991	-981	-971	-961	-951	-941	-931	-921
						GTTGACAGTT		mma a cocamorm
	CTAGGATATG	TTTTGGAATA						
				cmacoi Ge	nomic DNA			
				201	074	0.61	-851	-841
	-911	-901	-891	-88T	-8 /T	-861	-851	-041
	*	*	*					2002 2 2 2 000 M
	ATGTAAGAAA	CGACGAAATA				GAGCTAGATC	CIMANGAIRI	ATAAAAGTAT
				cmACOl Ge	nomic DNA_			>
						=0.4	550	500
	-831	-821	-811	-801	-791	-781	-771	-761
	*	*	*	*	*	*	*	
						TTCAAGGITA		
				cmACO1 Ge	nomic DNA_			>
	-751	-741	-731	-721	-711	-701	-691	-681
	*	*	*	*	*	*	*	*
	AAATTTCATA	GATAATGCAT				TGGTTGTCCT	AATTTTTGTA	GTAAATAAGC
				cmACO1 Ge	nomic DNA			>
-671	-661	-651	-641	-631	-621	-611	-601	
	*	*	*	*	*	*	*	*
	GTAGTTCAAG							CACGITIACGA
				cmACO1 Ge	nomic DNA			>

Fig. 1A

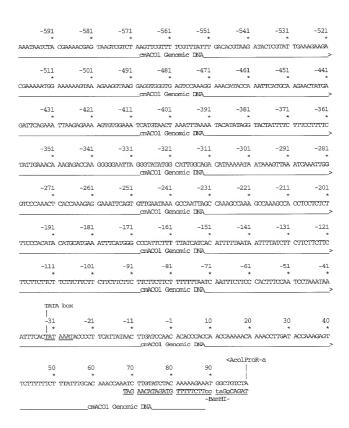


Fig. 1B

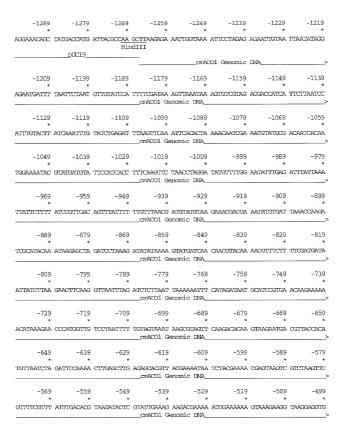


Fig. 2A

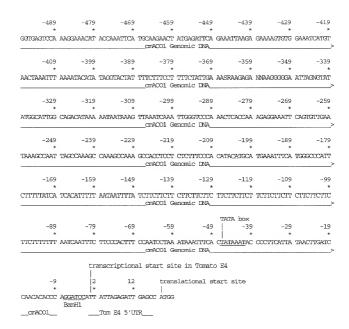


Fig. 2B

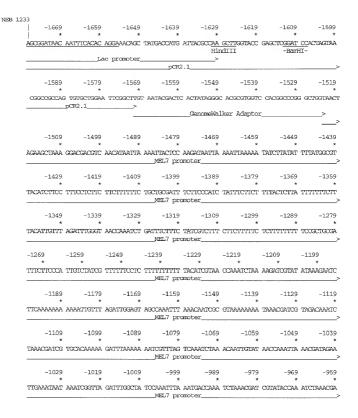


Fig. 3A

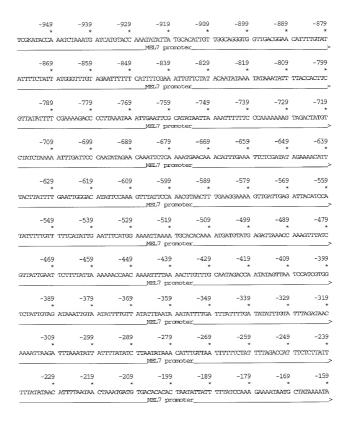


Fig. 3B

-143	-135	-129	-112	-103	- 22	-03	-15
*	*	*	*	*	*	*	*
TOGGTCTTCT	TTATCACCTT	CATGATAATT	ATGAAAAATA	AAATAAAATT	TAATTATATA	ATTCATTTCA	TCTAATCGTA
			MEL7 pro	moter			>
			TATA box	v			
-69	-59	-49			-19	-9	2
-09		-49	-39		-19	-9	4
*	*	*	*	*	*	*	*
CAAGCTAGAT	ATTACTATAT	CAACAACTTT	GTGTATAAAA	AGGGCAAGAA	ATTAAGCATT	ATCGTGTGAG	CCACTITITC
			MEL7 pro	moter			
		Me	17 translat	ional start	site		
		1		DruNcoSt			
		- 1	~0.00	J. Carocosc			
40							
12		32	42				
*	*	*	*	*			
TATATCTAGA	GATAGAAGGT	TTAAAATCAT	GTCTCTAATT	GGAAAGCTTG	TGAGT		
	TTCCA	AATTITGGTA	CCGAGACTAA	CCTTTCG			
		-Nco					
) ATT	. 7						
	L7 promoter						
			M	EL7 cds	>		

Fig. 3C

NEB 1233 TTGTGTGGAA TTGTGAGCCG ATAACAATTT CACACAGGAA ACAGCTATGA CCATGATTAC GCCAAGCTTG GTACCGAGCT Lac promoter____ -2068 -2058 -2048 -2038 -2028 -2018 -2008 * * * * * * * * * * * CGGATCCACT AGTAACGGCC GCCAGTGTGC TGGAATTCGG CTTACTATAG GGCACGCGTG GTCGACGGCC CGGGCTGGTC -SpeI--__pCR2.1__ GenomeWalker Adaptor____ -1998 -1988 -1978 -1968 -1958 -1948 -1938 -1928 * * * * * * * * * * CAATCACCGA ACATCATCTT ATCTAGGTGT CGGGAGATGC TACCTATCTG CTGATGTTGG TTTCTFTCTT TGAAAGATAC MEL2 promoter_ -1918 -1908 -1898 -1888 -1878 -1868 -1858 -1848 * * * * * * * * * * TCTCCTGACT TTTTAGTTGT TGCATCTAGA GATGTCCTCT ATTATTTTGA CACCTTTTCT TCTGACGGTG TAGAGCAACA MEL2 promoter_ CAAAAAAATC TIGAATTICT ATTAATGGAA TGAGCTATAT CIATACAAAT TGGAACCATA TGACAAATTA AGAAGATTCC MEL2 promoter_ TTTCTGAATA TTATGCAATA GAAATACTCA CCAGGTGTAA TGATGCACCT TATAGAGAAA ACTTCGACGA ACAAGAGACG ____MEL2 promoter_____ -1678 -1668 -1658 -1648 -1638 -1628 -1618 -1608 * * * * * * * * * * GCTACTAGGT TTTGGTGGAA TGGGTATTTC TGACCTACTA TGTTTCAGGA TGCGAGGATC TTCATGGTCA ATTGTGACCG ____MEL2 promoter____ -1598 -1588 -1578 -1568 -1558 -1548 -1538 -1528 * * * * * * * * * * * ATGGTGGAGA ACTGAAAATA TTTCCCATCT CAATGAAATA CTAAAACAAC ATATCTTAGA GGTTGAACTA TTTGATATCT ____MEL2 promoter____ -1518 -1508 -1498 -1488 -1478 -1468 -1458 -1448 * * * * * * * AGGGAATACA TITTATGGGA CCGITTICTA GTTGTTCCGG CAAACACGCA TTCGAGACGG GACGITCATG TCGCATACCA ____MEL2 promoter__

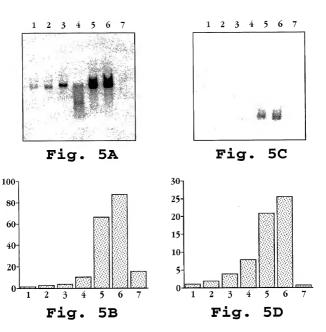
Fig. 4A

-1438	-1428	-1418	-1408	-1398	-1388	-1378	-1368
COGAGGATOC	GCATGTAAGC					* ATATATTTT	
			PESIZ DE	AIDCEL			
-1358 *	-1348	-1338	-1328	-1318 *	-1308 *	-1298 *	-1288
GATAGITICT	AAATTTGTTG			TCAGTCCTAG moter_		TOGGTACTTA	GCCACTCTCT >
-1278	-1268	-1258	-1248	-1238	-1228	-1218	-1208
*	*	*	*	*	*	*	*
GGGGACAAGA	TATGOGAGAT			actactcaaa moter		TGGGGACCTA	AG1C1AGGGC >
-1198	-1188	-1178			-1148	-1138	-1128
*	*	*	*		*	* CCAATTATGG	*
CLACAAGACG		GIAGITOCAC				CCAATTATIGG	ACTMAGCTIG
				andeer			
-1118	-1108	-1098	-1088	-1078	-1068	-1058 *	-1048 *
ATCAAGCTGT	GCAACGGATT	GAAGAACAAA	CAAGAAATCA	CGATGCGTTA	GCTTCAAAAG	TOGAATGAAT	GIGAAAGITC
			MEL2 pn	omoter			>
-1038	-1028	-1018	-1008	-998	-988	-978	-968
* 30030330303	messemercere	*	*	*	mmccccamacc	TATANNTATT	TO COMPANIE OF THE COMPANIE OF
MINGHONON	1GHG1CGGGC			romoter_			ICCATTALIC
-958	-948	-938	-928	-918	-908	-898	-888
*	*	*	*	*	*	*	*
TTAAGTTTTT	GAATTACAGT					ACTITIGIAT	AATTGTAGGA
			MEL2 pr	omoter			>
-878	-868	-858	_949	-838	-828	-818	-808
-676	-505	-050	-040	-030	-020	-010	-505
CCTGTGGTGT	AGAATGGCAT			TACGATTTTC		TTTTAACGAG	GAATATTTTT >
-798	-788	-778	-768	-758	-748	-738	-728
*	*	*	*	*	*	*	*
TATTTGTATT	ATGAACTTTA			TGTATTATGA omoter		TTTTGTTGAA	TTTTTGTTTG >
-718	-708	-698	-688	-678	-668	-658	-648
AATEMPPER	יוידעאַידיי	TAAATTTTAT	TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	TGAATCGATA	ACGAATGCAA	ATATTTTACG	TTYMAAAAAA

Fig. 4B

-638	-628	-618	-608	-598	-588	-578	-568
ATAGGAAAAT	ATTICAAAAA				CGACOCATTA		
			precia pro	andter			
-558 *	-548 *	-538 *	-528 *	-518	-508 *	-498 *	-488
CAAACATCAC	ATOGGGGATG	GTTATTACCG	ACOCATGAAT	GACACOGAAT	ATATAAACGT	AAGGAATAGT	TATTCCTGAC
			MEL2 pro	omoter			>
-478	-468 *	-458 *	-448 *	-438 *	-428 *	-418 *	-408 *
	TGTCGGAACT						
			MEL2 pro	omoter			>
-398 *	-388	-378 *	-368 *	-358 *	-348	-338	-328 *
GCTCCACTTC	TTGTAGTGAA						
							>
			niperre	ect inverte	d repeat		
-318	-308	-298	-288	-278	-268	-258	-248
-510	*	-250	*		*	*	*
ATAATCAATA	TMCAAAAGTT	CAATICCAAA	AATTACATTT	CTCTAGAAAT			AAAGGTTTTA
			MEL2 pro	amoter			>
-238	-228	-218	-208	100	-188	-178	-168
-230	-220	-210	-200	-130	-100	-1/0	-100
AGTGAATTGA	AAATTTCAAA	ACCITAATTICG	ATTRACCICAG	AAAATTATTT	TAATCACCAT	TCAAAAGTTA	TTTAACAATGA
~158	-148	-138	-128	-118	-108	-98	-88
*	*	*	*		*	*	*
AAAATATOGA	AGATAAGATT	TCAAAATTAC	GTAATTTACT	TCTACGTTTC	TITCTITCCC	CTTTAGTAAC	TTCACTCATA
			MEL2 pro	omoter			>
TATA b							
-78	-68	-58		-38	-28	-18	-8
*	*				*	*	*
TCTTTATATA	CGITCCATCC						ACTCACCCIT
			MEL2 pr	amoter			>
	2 translati		site				
	<mel2_< td=""><td></td><td></td><td></td><td></td><td></td><td></td></mel2_<>						
1 3	13						
		1 1 1 1					
	GAAACAATGC						
-NcoI	_CTTTGTTACG	1116					
pro							
	MET.2 cds						

Fig. 4C



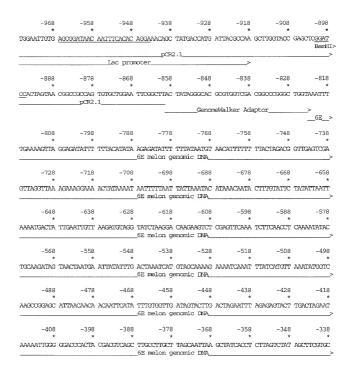


Fig. 6A

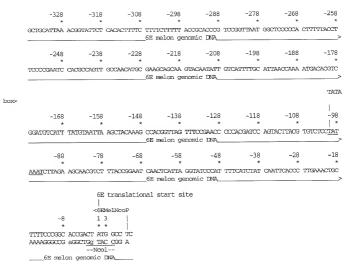


Fig. 6B

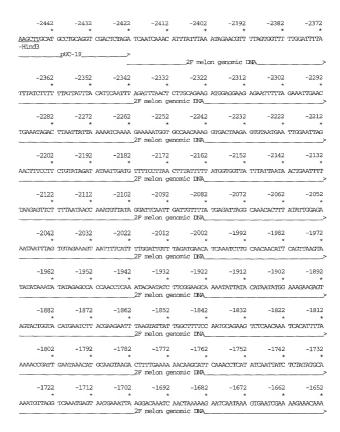


Fig. 7A

-1572	-1582	-1592	-1602	-1612	-1622	-1632	-1642
2202220000	TTTGATAAAC	*	~	*	*	*	*
APLAPPICIC >				ATGATGCAGT 2F melon de		ACCIAAIGIA	TATCAATCAA
			nonce brag	zr neion ge			
-1492	-1502	-1512	-1522	-1532	-1542	-1552	-1562
*	*	*	*	*	*	*	*
	GTCCAAAACA					CGAGAAGAGA	CACTOCAATA
>			nomic DNA	2F melon g∈			
-1412	-1422	-1432	-1442	-1452	-1462	-1472	-1482
-1412	-1422	-1432	-1442	-1452	-1462	-14/2	-1482
CITCTTTCAA	TGTGAGTTTT	GTACATGTGG	AGACATGGTT	TGGCAGGGAA	ТАТСААТААА	GAAGAAATGT	AAAACATGGT
>				2F melon qe		G12411101	IEEE KAILOOI
				-			
-1332	-1342	-1352	-1362	-1372	-1382	-1392	-1402
*	*	*	*	*	*	*	*
TGGCGIGAAI	GGCTACCCCA	ACCATGACAG	CIGITIGGAA	AAGAGAAACA	AOGACCCAAC	AAATTGGATT	ATCTGTGAAT
)T—-			nomia DNA	2F melon de			
			HOME DAY	_zr meton ge			
-1252	-1262	-1272	-1282	-1292	-1302	-1312	-1322
*	*	*	*	*	*	*	*
TATATACGAA	TCTTATATTT	TCGATTCATA	GTTTTTTTAT	CCCGCCATTC	AGCTCTCATC	TAATTAATTA	ATCAAGTATT
>			enomic DNA_	2F melon ge			
-1172	-1182	-1192	-1202	-1212	-1222	-1232	-1242
ma a amerecana	ATATAAAAAG	CATING CATALOG S'GL	*	· · · · · · · · · · · · · · · · · · ·	*	*	*
TARATOCOMA				2F melon of			TAATTCTTGA
			TORRIC DAY	_zr neion ge			
-1092	-1102	-1112	-1122	-1132	-1142	-1152	-1162
*	*	*	*	*	*	*	*
	AGGGAAATGG					AGTATTCACT	TGATTCTATT
>			enomic DNA_	_2F melon go			
1010	1000	4000					
-1012	-1022	-1032	-1042	-1052	-1062	-1072	-1082
מ מיצווווווווווווווווווווווווווווווווווו	AGTICATOCTC	A CONTRACTOR OF THE ACTION OF	macomma oum	*	*	~	w
MIIIIICAN				2F melon q		GITAAATTIC	GIGATIGGIA
			monte man_	_zr neion g			
-932	-942	-952	-962	-972	-982	-992	-1002
*	*	*	*	*	*	*	*
	GAAAAGAAGT					CCAATTATAC	CICICAACIT
>			enomic DNA_	_2F melon g			
050	0.50	000					
-852	-862	-872	-882	-892	-902	-912	-922
AATCAAGTTG	THE THE PERSON IN THE PERSON I	minimization of the	ריים מיוימיוידים	WILLIAM WILLIA	manaca com	meanmeane	»
AMICAMOTIO						I CHAI I CAAC	ACTITITIT
			Transco Trans	_nrion 9			

Fig. 7B

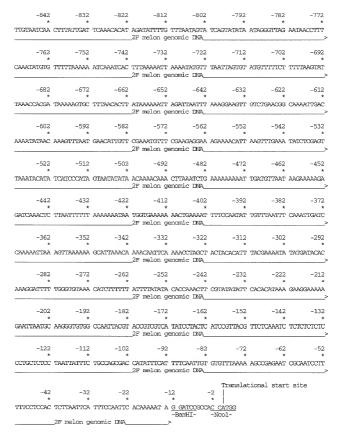


Fig. 7C

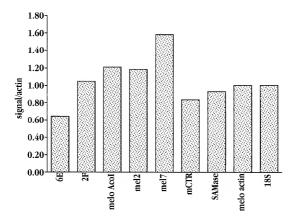


Fig. 8